

## RAW SEQUENCE LISTING

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Application Serial Number: 10/573,821 A  
Source: IFVO  
Date Processed by STIC: 8/22/06

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IFWO

## RAW SEQUENCE LISTING

DATE: 08/22/2006

PATENT APPLICATION: US/10/573,821A

TIME: 10:43:06

Input Set : A:\47259501.APP

Output Set: N:\CRF4\08222006\J573821A.raw

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3 <110> APPLICANT: OKUNO, KAZUAKI
4     YABUTA, MASAYUKI
6 <120> TITLE OF INVENTION: POLYPEPTIDE CLEAVAGE METHOD USING OMPT PROTEASE VARIANT
8 <130> FILE REFERENCE: 47259.5001/00US
10 <140> CURRENT APPLICATION NUMBER: 10/573,821A
11 <141> CURRENT FILING DATE: 2006-03-28
13 <150> PRIOR APPLICATION NUMBER: PCT/JP04/014704
14 <151> PRIOR FILING DATE: 2004-09-29
16 <150> PRIOR APPLICATION NUMBER: JP 2003-342183
17 <151> PRIOR FILING DATE: 2003-09-30
19 <160> NUMBER OF SEQ ID NOS: 38
21 <170> SOFTWARE: PatentIn Ver. 3.3
23 <210> SEQ ID NO: 1
24 <211> LENGTH: 184
25 <212> TYPE: PRT
26 <213> ORGANISM: Artificial Sequence
28 <220> FEATURE:
29 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
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36 Trp Glu Asn Pro Gly Val Thr Gln Leu Asn Arg Leu Ala Ala His Pro
37             20             25             30
39 Pro Phe Ala Ser Trp Arg Asn Ser Asp Asp Ala Arg Thr Asp Arg Pro
40             35             40             45
42 Ser Gln Gln Leu Arg Ser Leu Asn Gly Glu Trp Arg Phe Ala Trp Phe
43   50             55             60
45 Pro Ala Pro Glu Ala Val Pro Glu Ser Leu Leu Asp Leu Pro Glu Ala
46   65             70             75             80
48 Asp Thr Val Val Val Pro Asp Ser Ser Asn Trp Gln Met His Gly Tyr
49             85             90             95
51 Asp Ala Pro Ile Tyr Thr Asn Val Thr Tyr Pro Ile Thr Val Asn Pro
52             100            105            110
54 Pro Phe Val Pro Thr Glu Pro His His His His Pro Gly Gly Arg Gln
55             115            120            125
57 Met His Gly Tyr Asp Ala Glu Leu Arg Leu Tyr Arg Arg His His Gly
58   130            135            140
60 Ser Gly Ser Pro Tyr Arg His Pro Arg His Ala Glu Gly Thr Phe Thr
61 145            150            155            160
63 Ser Asp Val Ser Ser Tyr Leu Glu Gly Gln Ala Ala Lys Glu Phe Ile
64             165            170            175
66 Ala Trp Leu Val Lys Gly Arg Gly

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84          20          25          30
86 Pro Phe Ala Ser Trp Arg Asn Ser Asp Asp Ala Arg Thr Asp Arg Pro
87          35          40          45
89 Ser Gln Gln Leu Arg Ser Leu Asn Gly Glu Trp Arg Phe Ala Trp Phe
90          50          55          60
92 Pro Ala Pro Glu Ala Val Pro Glu Ser Leu Leu Asp Leu Pro Glu Ala
93   65          70          75          80
95 Asp Thr Val Val Val Pro Asp Ser Ser Asn Trp Gln Met His Gly Tyr
96          85          90          95
98 Asp Ala Pro Ile Tyr Thr Asn Val Thr Tyr Pro Ile Thr Val Asn Pro
99          100          105          110
101 Pro Phe Val Pro Thr Glu Pro His His His His Pro Gly Gly Arg Gln
102          115          120          125
104 Met His Ala Ala Ala Ala Ala Ala Ala Ala Arg Arg Ala Ala Ala
105          130          135          140
107 Ala Gly Ser Pro Tyr Arg His Pro Arg His Ala Glu Gly Thr Phe Thr
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131          20          25          30
133 Pro Phe Ala Ser Trp Arg Asn Ser Asp Asp Ala Arg Thr Asp Arg Pro
134          35          40          45
136 Ser Gln Gln Leu Arg Ser Leu Asn Gly Glu Trp Arg Phe Ala Trp Phe
137          50          55          60

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139 Pro Ala Pro Glu Ala Val Pro Glu Ser Leu Leu Asp Leu Pro Glu Ala
140 65 70 75 80
142 Asp Thr Val Val Val Pro Asp Ser Ser Asn Trp Gln Met His Gly Tyr
143 85 90 95
145 Asp Ala Pro Ile Tyr Thr Asn Val Thr Tyr Pro Ile Thr Val Asn Pro
146 100 105 110
148 Pro Phe Val Pro Thr Glu Pro His His His His Pro Gly Gly Arg Gln
149 115 120 125
151 Met His Ala Ala Ala Ala Ala Ala Ala Ala Arg Arg Ala Arg Ala
152 130 135 140
154 Ala Gly Ser Pro Tyr Arg His Pro Arg His Ala Glu Gly Thr Phe Thr
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178 20 25 30
180 Pro Phe Ala Ser Trp Arg Asn Ser Asp Asp Ala Arg Thr Asp Arg Pro
181 35 40 45
183 Ser Gln Gln Leu Arg Ser Leu Asn Gly Glu Trp Arg Phe Ala Trp Phe
184 50 55 60
186 Pro Ala Pro Glu Ala Val Pro Glu Ser Leu Leu Asp Leu Pro Glu Ala
187 65 70 75 80
189 Asp Thr Val Val Val Pro Asp Ser Ser Asn Trp Gln Met His Gly Tyr
190 85 90 95
192 Asp Ala Pro Ile Tyr Thr Asn Val Thr Tyr Pro Ile Thr Val Asn Pro
193 100 105 110
195 Pro Phe Val Pro Thr Glu Pro His His His His Pro Gly Gly Arg Gln
196 115 120 125
198 Met His Ala Ala Ala Ala Ala Ala Ala Ala Arg Arg Arg Ala Arg Ala
199 130 135 140
201 Ala Gly Ser Pro Tyr Arg His Pro Arg His Ala Glu Gly Thr Phe Thr
202 145 150 155 160
204 Ser Asp Val Ser Ser Tyr Leu Glu Gly Gln Ala Ala Lys Glu Phe Ile
205 165 170 175
207 Ala Trp Leu Val Lys Gly Arg Gly
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211 <210> SEQ ID NO: 5

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218 protein sequence

220 &lt;400&gt; SEQUENCE: 5

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225           20           25           30
227 Pro Phe Ala Ser Trp Arg Asn Ser Asp Asp Ala Arg Thr Asp Arg Pro
228           35           40           45
230 Ser Gln Gln Leu Arg Ser Leu Asn Gly Glu Trp Arg Phe Ala Trp Phe
231           50           55           60
233 Pro Ala Pro Glu Ala Val Pro Glu Ser Leu Leu Asp Leu Pro Glu Ala
234   65           70           75           80
236 Asp Thr Val Val Val Pro Asp Ser Ser Asn Trp Gln Met His Gly Tyr
237           85           90           95
239 Asp Ala Pro Ile Tyr Thr Asn Val Thr Tyr Pro Ile Thr Val Asn Pro
240           100          105          110
242 Pro Phe Val Pro Thr Glu Pro His His His His Pro Gly Gly Arg Gln
243           115          120          125
245 Met His Gly Tyr Asp Ala Glu Leu Arg Leu Tyr Arg Phe Val Pro Ile
246           130          135          140
248 Phe Thr Tyr Gly Glu Leu Gln Arg Met Gln Glu Lys Glu Arg Asn Lys
249 145          150          155          160
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261 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic  
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264 &lt;400&gt; SEQUENCE: 6

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268 Trp Glu Asn Pro Gly Val Thr Gln Leu Asn Arg Leu Ala Ala His Pro
269           20           25           30
271 Pro Phe Ala Ser Trp Arg Asn Ser Asp Asp Ala Arg Thr Asp Arg Pro
272           35           40           45
274 Ser Gln Gln Leu Arg Ser Leu Asn Gly Glu Trp Arg Phe Ala Trp Phe
275           50           55           60
277 Pro Ala Pro Glu Ala Val Pro Glu Ser Leu Leu Asp Leu Pro Glu Ala
278   65           70           75           80
280 Asp Thr Val Val Val Pro Asp Ser Ser Asn Trp Gln Met His Gly Tyr
281           85           90           95
283 Asp Ala Pro Ile Tyr Thr Asn Val Thr Tyr Pro Ile Thr Val Asn Pro

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284          100          105          110
286 Pro Phe Val Pro Thr Glu Pro His His His His Pro Gly Gly Arg Gln
287          115          120          125
289 Met His Ala Ala Ala Ala Ala Ala Ala Arg Arg Arg Ala Arg Phe
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313          20          25          30
315 Pro Phe Ala Ser Trp Arg Asn Ser Asp Asp Ala Arg Thr Asp Arg Pro
316          35          40          45
318 Ser Gln Gln Leu Arg Ser Leu Asn Gly Glu Trp Arg Phe Ala Trp Phe
319          50          55          60
321 Pro Ala Pro Glu Ala Val Pro Glu Ser Leu Leu Asp Leu Pro Glu Ala
322 65          70          75          80
324 Asp Thr Val Val Val Pro Asp Ser Ser Asn Trp Gln Met His Gly Tyr
325          85          90          95
327 Asp Ala Pro Ile Tyr Thr Asn Val Thr Tyr Pro Ile Thr Val Asn Pro
328          100          105          110
330 Pro Phe Val Pro Thr Glu Pro His His His His Pro Gly Gly Arg Gln
331          115          120          125
333 Met His Ala Ala Ala Ala Ala Ala Ala Arg Arg Arg Ala Arg Ser
334          130          135          140
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339 Arg Pro Val Lys Val Tyr Pro
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**VERIFICATION SUMMARY**

DATE: 08/22/2006

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